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<110> SHITARA, KENYA SHIBUYA, MASABUMI												
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gaa tgg att gg Glu Trp Ile Gl					240
cag aag ttc aa Gln Lys Phe Ly 6	s Gly Lys				288

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ccc aaa ctc ttg att Pro Lys Leu Leu Ile 45			55 5	240
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gtc tat gcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag
                                                                   96
Val Tyr Ala Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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         -1
cct ggc gcc tca gtg aag gtc tcc tgc aag gct tct gga tac acc ttc
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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age agt aac tat ata agt tgg gtg cga cag gee eet gga caa ggg ett
                                                                   192
Ser Ser Asn Tyr Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
30
gag tgg atg gga tgg att tat gct gga act ggt gat gcc agc tat aat
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Glu Trp Met Gly Trp Ile Tyr Ala Gly Thr Gly Asp Ala Ser Tyr Asn
                 50
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Gln Lys Phe Thr Ala Arg Val Thr Ile Thr Val Asp Thr Ser Thr Ser
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             65
                                                                   336
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Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
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                             85
                                                  90
tat tac tgt gcg aga cac ggg ggg gac ggc tac tgg ttt gct tac tgg
                                                                   384
Tyr Tyr Cys Ala Arg His Gly Gly Asp Gly Tyr Trp Phe Ala Tyr Trp
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                                                                   415
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gtc cac tcc cag gtg cag ctg gtg cag tcc gga gct gag gtg aag aag
                                                                   96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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              1
cct ggg gcc tca gtg aag gtc tcc tgc aag gct tct gga tac acc ttc
                                                                   144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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Ile Asn Tyr Asn Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
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                     35
gag tgg atg gga gct att ttt cca gga aat ggt ttt act tcc tac aat
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Glu Trp Met Gly Ala Ile Phe Pro Gly Asn Gly Phe Thr Ser Tyr Asn
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Gln Lys Phe Lys Gly Arg Val Thr Ile Thr Val Asp Lys Ser Thr Ser
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aca gcc tac atg gag ctg agc ctg aga tct gag gac acg gcc gtg
Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
                             85
tat tac tgt gcg aga gat ggt gac tat tac ttt gac tac tgg ggc cag
Tyr Tyr Cys Ala Arg Asp Gly Asp Tyr Tyr Phe Asp Tyr Trp Gly Gln
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gga acc ctg gtc acc gtc tcc tca g
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gtc ata ata tcc aga gga gat atc cag atg acc cag tct cca tcc tcc
                                                                   96
Val Ile Ile Ser Arg Gly Asp Ile Gln Met Thr Gln Ser Pro Ser Ser
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ctg tct gca tct gta gga gac aga gtc acc atc act tgt agt gct agc
                                                                   144
Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Ser Ala Ser
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tca agt gtc agt tac atg cac tgg tat cag cag aaa cca ggg aaa gcc
Ser Ser Val Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Lys Ala
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                                                      40
cct aag ctt ctg atc tat gac aca tcc aaa ctg cct tct ggg gtc cca
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Pro Lys Leu Leu Ile Tyr Asp Thr Ser Lys Leu Pro Ser Gly Val Pro
         45
                                                                   288
tca agg ttc agc ggc agt gga tct ggg aca gat ttc act ctc acc atc
Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile
    60
                                                                   336
age age etg cag cet gaa gat tit gea act tat tae tgt cag cag tgg
Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Trp
75
agt agt aac cca ccc acg ttc ggc caa ggg acc aag gta gag atc aaa c 385
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50

45

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gac ega tte agt gge age ggg tet ggg aca gat tte act etc ace ate
                                                                   288
Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile
age age etg cag get gaa gae gte gea gtt tat tae tgt eat eag tgg
Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys His Gln Trp
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agt atg tac acg ttc ggc caa ggg acc aag gtg gag atc aaa c
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Ser Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys
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<220> <223> Description of Artificial Sequence: Synthetic DNA	
<400> 60 cataaatcct ataggtacca acgacaacta	30
<210> 61 <211> 87 <212> DNA <213> Artificial Sequence	

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<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 61
caggaaacag ctatgacgaa ttccaccatg gattttcaag tgcagatttt cagcttcctg 60
ctaatcagtg cctcagtcat aatatcc
<210> 62
<211> 93
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<400> 62
aagtgatggt gactctgtct cctacagatg cagacaggga ggatggagac tgggtcatct 60
ggatatetee tetggatatt atgactgagg cac
<210> 63
<211> 85
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<400> 63
agacagagtc accatcactt gtagtgccag ctcgagtgta agttacatgc actggtatca 60
gcagaaacca gggaaagccc ctaag
<210> 64
<211> 84
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<400> 64
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gaagettagg ggettteeet ggtt
<210> 65
<211> 94
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 65
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tgaagatttt gcaacttatt actgtcatca gtgg
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<210> 66
<211> 85
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<213> Artificial Sequence
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<400> 66
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atactccact gatgacagta ataag
<210> 67
<211> 379
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<221> sig_peptide
<222> (1)..(66)
<220>
<221> mat_peptide
<222> (67)..(378)
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<221> CDS
<222> (1)..(378)
<400> 67
atg gat ttt caa gtg cag att ttc agc ttc ctg cta atc agt gcc tca
Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser
                            -15
gtc ata ata tcc aga gga gat atc cag atg acc cag tct cca tcc tcc
Val Ile Ile Ser Arg Gly Asp Ile Gln Met Thr Gln Ser Pro Ser Ser
ctg tct gca tct gta gga gac aga gtc acc atc act tgt agt gcc agc
Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Ser Ala Ser
                 15
                                     20
tcg agt gta agt tac atg cac tgg tat cag cag aaa cca ggg aaa gcc
Ser Ser Val Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Lys Ala
cct aag ctt ctg atc tat aga aca tcc aac ctg gct tct ggg gtc cca
Pro Lys Leu Leu Ile Tyr Arg Thr Ser Asn Leu Ala Ser Gly Val Pro
         45
                             50
```

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```
tca agg ttc agc ggc agt gga tct ggg aca gat ttc act ctc acc atc
                                                                   288
Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile
     60
                         65
age age etg cag eet gaa gat tit gea aet tat tae tgt eat eag tgg
                                                                   336
Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys His Gln Trp
                     80
                                         85
agt atg tac acg ttc ggc caa ggg acc aag gta gag atc aaa c
                                                                   379
Ser Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys
                 95
<210> 68
<211> 80
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
caggaaacag ctatgactcc ggagctgagg tgaagaagcc tggggcctca gtgaaggtct 60
cctgcaaggc ttctggatac
<210> 69
<211> 80
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 69
ccactcaagc ccttgtccag gggcctgtcg cacccagtgc atattgtaat taatgaaggt 60
gtatccagaa gccttgcagg
<210> 70
<211> 81
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
ctggacaagg gcttgagtgg atgggagcta tttttccagg aaatggtttt acttcctaca 60
atcagaagtt caagggcaga g
<210> 71
<211> 79
<212> DNA
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<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Synthetic DNA
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teteaggetg egeagetgea tgtaggetgt getegtggae ttgtegaegg taatggtgae 60
tetgeeettg aacttetga
<210> 72
<211> 83
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<400> 72
tgcagctgcg cagcctgaga tctgaggaca cggccgtgta tttctgtgcg agagatggtg 60
actattactt tgactactgg ggc
<210> 73
<211> 81
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<400> 73
gttttcccag tcacgacggg cccttggtgg aggctgagga gacggtgacc agggttccct 60
ggccccagta gtcaaagtaa t
<210> 74
<211> 409
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<220>
<221> sig_peptide
<222> (1)..(57)
<220>
<221> mat_peptide
<222> (58)..(408)
<220>
<221> CDS
<222> (1)..(408)
<400> 74
atg gga ttc agc agg atc ttt ctc ttc ctc ctg tca gtg act aca ggt
                                                                   48
Met Gly Phe Ser Arg Ile Phe Leu Phe Leu Leu Ser Val Thr Thr Gly
                                     -10
```

												gag Glu 10				96
												gga Gly				144
												gga Gly				192
												act Thr				240
												aag Lys				288
												gac Asp 90				336
		_		_	_		_				_	tac Tyr			_	384
					gtc Val 115			g								409
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<220 <223		escri	iptio	on of	Art	ific	cial	Sequ	ience	e: S	nthe	etic	DNA			
cago		ag d			ia tt it aa			g gat	tttc	caag	tgca	ıgatt	tt c	agct	tectg	60 87
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<220> <223> Description of Artificial Sequence: Synthetic DNA																
aagt		gt c			t co					gga	ggat	ggag	jac t	gggt	catct	60 93

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<210> 77
<211> 85
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<213> Artificial Sequence
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agaggaggtc accatcactt gtagtgccag ctcgagtgta agttacatgc actggtatca 60
gcagaaacca gggaaagccc ctaag
<210> 78
<211> 84
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<400> 78
atccactgcc gctgaacctt gatgggaccc cagaagccag gttggatgtt ctatagatca 60
gaagettagg ggettteeet ggtt
<210> 79
<211> 94
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<400> 79
aaggttcagc ggcagtggat ctgggacatt ttatactctc accatcagca gcctgcagcc 60
tgaagatttt gcaacttatt actgtcatca gtgg
<210> 80
<211> 85
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<400> 80
gttttcccag tcacgaccgt acgtttgatc tctaccttgg tcccttggcc gaacgtgtac 60
atactccact gatgacagta ataag
<210> 81
<211> 379
<212> DNA
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<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Synthetic DNA
<221> sig_peptide
<222> (1)..(66)
<220>
<221> mat_peptide
<222> (67)..(378)
<220>
<221> CDS
<222> (1)..(378)
<400> 81
atg gat ttt caa gtg cag att ttc agc ttc ctg cta atc agt gcc tca
Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser
gtc ata ata tcc aga gga gat atc cag atg acc cag tct cca tcc tcc
Val Ile Ile Ser Arg Gly Asp Ile Gln Met Thr Gln Ser Pro Ser Ser
ctg tct gca tct gta gga gag gtc acc atc act tgt agt gcc agc
                                                                   144
Leu Ser Ala Ser Val Gly Glu Glu Val Thr Ile Thr Cys Ser Ala Ser
tcg agt gta agt tac atg cac tgg tat cag cag aaa cca ggg aaa gcc
                                                                   192
Ser Ser Val Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Lys Ala
                                 35
cct aag ctt ctg atc tat aga aca tcc aac ctg gct tct ggg gtc cca
                                                                   240
Pro Lys Leu Leu Ile Tyr Arg Thr Ser Asn Leu Ala Ser Gly Val Pro
        45
tca agg ttc agc ggc agt gga tct ggg aca ttt tat act ctc acc atc
Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Phe Tyr Thr Leu Thr Ile
    60
                         65
age age etg cag eet gaa gat tit gea aet tat tae tgt eat eag tgg
Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys His Gln Trp
75
                     80
                                         85
                                                                   379
agt atg tac acg ttc ggc caa ggg acc aag gta gag atc aaa c
Ser Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys
                 95
                                    100
<210> 82
<211> 138
<212> PRT
<213> Mus musculus
Met Glu Trp Asn Trp Val Val Leu Phe Leu Leu Ser Leu Thr Ala Gly
                -15
                                    -10
```

Val Tyr Ala Gln Gly Gln Met Gln Gln Ser Gly Ala Glu Leu Val Lys
-1 1 5 10

Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Pro Ser Gly Phe Thr Phe 15 20 25

Ser Ser Asn Tyr Ile Ser Trp Leu Lys Gln Lys Pro Gly Gln Ser Leu 30 35 40 45

Glu Trp Ile Ala Trp Ile Tyr Ala Gly Thr Gly Asp Ala Ser Tyr Asn
50 55 60

Gln Lys Phe Thr Ala Lys Ala His Val Thr Val Asp Thr Ser Ser Ser 65 70 75

Thr Ala Tyr Met Gln Leu Ser Ser Leu Thr Thr Glu Asp Ser Ala Ile 80 85 90

Tyr Tyr Cys Ala Arg His Gly Gly Asp Gly Tyr Trp Phe Ala Tyr Trp 95 100 105

Gly Gln Gly Thr Leu Val Thr Val Ser Ala 110 115

<210> 83

<211> 128

<212> PRT

<213> Mus musculus

<400> 83

Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser
-20 -15 -10

Val Ile Ile Ser Arg Gly Gln Leu Val Leu Thr Gln Ser Pro Ala Ile
-5 -1 1 5 10

Met Ser Ala Ser Gln Gly Glu Lys Val Thr Met Thr Cys Ser Ala Ser 15 20 25

Ser Ser Val Ser Tyr Met His Trp Tyr Gln Gln Lys Ser Gly Thr Ser 30 35 40

Pro Lys Arg Trp Ile Tyr Asp Thr Ser Lys Leu Pro Ser Gly Val Pro
45 50 55

Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile
60 65 70

Ser Ser Met Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Trp
75 80 85 90

Ser Ser Asn Pro Pro Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys 95 100 105 <211> 136

<212> PRT

<213> Mus musculus

<400> 84

Met Gly Phe Ser Arg Ile Phe Leu Phe Leu Leu Ser Val Thr Thr Gly
-15 -10 -5

Val His Ser Gln Ala Phe Leu Gln Gln Ser Gly Ala Glu Leu Val Arg

Pro Gly Ala Ser Val Lys Met Ser Cys Lys Ala Ser Gly Tyr Thr Phe 15 20 25

Ile Asn Tyr Asn Met His Trp Val Lys Gln Thr Pro Arg Gln Gly Leu 30 35 40 45

Glu Trp Ile Gly Ala Ile Phe Pro Gly Asn Gly Phe Thr Ser Tyr Asn 50 $$ 55 $$ 60

Gln Lys Phe Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser 65 70 75

Thr Val Tyr Met Gln Leu Arg Ser Leu Thr Ser Glu Asp Ser Ala Val 80 85 90

Tyr Phe Cys Ala Arg Asp Gly Asp Tyr Tyr Phe Asp Tyr Trp Gly Gln 95 100 105

Gly Thr Thr Leu Thr Val Ser Ser 110 115

<210> 85

<211> 126

<212> PRT

<213> Mus musculus

<400> 85

Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser

Val Ile Met Ser Arg Gly Gln Ile Val Leu Thr Gln Ser Pro Ala Ile
-5 -1 1 5 10

Met Ser Ala Ser Leu Gly Glu Glu Ile Thr Leu Thr Cys Ser Ala Ser 15 20 25

Ser Ser Val Ser Tyr Met His Trp Tyr Gln Gln Lys Ser Gly Thr Ser 30 35 40

Pro Lys Leu Leu Ile Tyr Arg Thr Ser Asn Leu Ala Ser Gly Val Pro
45 50 55

Phe Arg Phe Ser Gly Ser Gly Ser Gly Thr Phe Tyr Ser Leu Thr Ile 60 65 70 Ser Ser Val Glu Ala Glu Asp Ala Ala Asp Tyr Tyr Cys His Gln Trp
75 80 85 90

Ser Met Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys 95 100

<210> 86

<211> 138

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic humanized antibody

<400> 86

Met Glu Trp Asn Trp Val Val Leu Phe Leu Leu Ser Leu Thr Ala Gly
-15 -10 -5

Val Tyr Ala Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
-1 1 5 10

Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
15 20 25

Ser Ser Asn Tyr Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu 30 40 45

Glu Trp Met Gly Trp Ile Tyr Ala Gly Thr Gly Asp Ala Ser Tyr Asn
50 55 60

Gln Lys Phe Thr Ala Arg Val Thr Ile Thr Val Asp Thr Ser Thr Ser 65 70 75

Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val 80 85 90

Tyr Tyr Cys Ala Arg His Gly Gly Asp Gly Tyr Trp Phe Ala Tyr Trp 95 100 105

Gly Gln Gly Thr Leu Val Thr Val Ser Ser 110 115

<210> 87

<211> 136

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic humanized antibody

<400> 87

Met Gly Phe Ser Arg Ile Phe Leu Phe Leu Leu Ser Val Thr Thr Gly

Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
-1 1 5 10

Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe 15 20 25

Ile Asn Tyr Asn Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu 30 35 40 45

Glu Trp Met Gly Ala Ile Phe Pro Gly Asn Gly Phe Thr Ser Tyr Asn 50 55 60

Gln Lys Phe Lys Gly Arg Val Thr Ile Thr Val Asp Lys Ser Thr Ser 65 70 75

Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val 80 85 90

Tyr Tyr Cys Ala Arg Asp Gly Asp Tyr Tyr Phe Asp Tyr Trp Gly Gln 95 100 105

Gly Thr Leu Val Thr Val Ser Ser 110 115

<210> 88

<211> 128

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic humanized antibody

<400> 88

Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser
-20 -15 -10

Val Ile Ile Ser Arg Gly Asp Ile Gln Met Thr Gln Ser Pro Ser Ser
-5 -1 1 5 10

Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Ser Ala Ser 15 20 25

Ser Ser Val Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Lys Ala 30 35 40

Pro Lys Leu Leu Ile Tyr Asp Thr Ser Lys Leu Pro Ser Gly Val Pro
45 50 55

Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile 60 65 70

Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Trp
75 80 85 90

Ser Ser Asn Pro Pro Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys 95 100 105

<210> 89

<211> 126

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic humanized antibody

<400> 89

Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser
-20 -15 -10

Val Ile Met Ser Arg Gly Asp Ile Val Met Thr Gln Ser Pro Asp Ser
-5 -1 1 5 10

Leu Ala Val Ser Leu Gly Glu Arg Ala Thr Ile Asn Cys Ser Ala Ser 15 20 25

Ser Ser Val Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Gln Pro $30 \ \ 35 \ \ 40$

Pro Lys Leu Leu Ile Tyr Arg Thr Ser Asn Leu Ala Ser Gly Val Pro
45 50 55

Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile 60 65 70

Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys His Gln Trp
75 80 85 90

Ser Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys 95 100

<210> 90

<211> 126

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic humanized antibody

<400> 90

Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser
-20 -15 -10

Val Ile Ile Ser Arg Gly Asp Ile Gln Met Thr Gln Ser Pro Ser Ser
-5 -1 1 5 10

Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Ser Ala Ser
15 20 25

Ser Ser Val Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Lys Ala 30 35 40

Pro Lys Leu Leu Ile Tyr Arg Thr Ser Asn Leu Ala Ser Gly Val Pro
45 50 55

Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile 60 65 70

Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys His Gln Trp 75 80 85 90

Ser Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys 95 100

<210> 91

<211> 136

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic humanized antibody

<400> 91

Met Gly Phe Ser Arg Ile Phe Leu Phe Leu Leu Ser Val Thr Thr Gly

Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
-1 1 5 10

Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe 15 20 25

Ile Asn Tyr Asn Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu 30 40 45

Glu Trp Met Gly Ala Ile Phe Pro Gly Asn Gly Phe Thr Ser Tyr Asn
50 55 60

Gln Lys Phe Lys Gly Arg Val Thr Ile Thr Val Asp Lys Ser Thr Ser 65 70 75

Thr Ala Tyr Met Gln Leu Arg Ser Leu Arg Ser Glu Asp Thr Ala Val 80 85 90

Tyr Phe Cys Ala Arg Asp Gly Asp Tyr Tyr Phe Asp Tyr Trp Gly Gln 95 100 105

Gly Thr Leu Val Thr Val Ser Ser 110 115

<210> 92

<211> 126

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic humanized antibody

<400> 92

Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser
-20 -15 -10

Val Ile Ile Ser Arg Gly Asp Ile Gln Met Thr Gln Ser Pro Ser Ser
-5 -1 1 5 10

Leu Ser Ala Ser Val Gly Glu Val Thr Ile Thr Cys Ser Ala Ser
15 20 25

Ser Ser Val Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Lys Ala 30 35 40

Pro Lys Leu Leu Ile Tyr Arg Thr Ser Asn Leu Ala Ser Gly Val Pro
45 50 55

Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Phe Tyr Thr Leu Thr Ile 60 65 70

Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys His Gln Trp 75 80 85 90

Ser Met Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys 95 100

<210> 93

<211> 117

<212> PRT

<213> Mus musculus

<400> 93

Gln Ala Phe Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ala 1 5 10 15

Ser Val Lys Met Ser Cys Lys Ala Ser Gly Tyr Thr Phe Ile Asn Tyr
20 25 30

Asn Met His Trp Val Lys Gln Thr Pro Arg Gln Gly Leu Glu Trp Ile 35 40 45

Gly Ala Ile Phe Pro Gly Asn Gly Phe Thr Ser Tyr Asn Gln Lys Phe 50 55 60

Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Val Tyr
65 70 75 80

Met Gln Leu Arg Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe Cys 85 90 95

Ala Arg Asp Gly Asp Tyr Tyr Phe Asp Tyr Trp Gly Gln Gly Thr Thr 100 105 110 Leu Thr Val Ser Ser 115

<210> 94

<211> 117

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic humanized antibody

<400> 94

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala 1 5 10 15

Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Ile Asn Tyr 20 25 30

Asn Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met 35 40 45

Gly Ala Ile Phe Pro Gly Asn Gly Phe Thr Ser Tyr Asn Gln Lys Phe 50 60

Lys Gly Arg Val Thr Ile Thr Val Asp Lys Ser Thr Ser Thr Ala Tyr 65 70 75 80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Arg Asp Gly Asp Tyr Tyr Phe Asp Tyr Trp Gly Gln Gly Thr Leu 100 105 110

Val Thr Val Ser Ser 115

<210> 95

<211> 117

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
humanized antibody

<400> 95

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala 1 5 10

Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Ile Asn Tyr
20 25 30

Asn Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met 35 40 45 Gly Ala Ile Phe Pro Gly Asn Gly Phe Thr Ser Tyr Asn Gln Lys Phe 50 55 60

Lys Gly Arg Val Thr Ile Thr Val Asp Lys Ser Thr Ser Thr Ala Tyr 65 70 75 80

Met Gln Leu Arg Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Phe Cys
85 90 95

Ala Arg Asp Gly Asp Tyr Tyr Phe Asp Tyr Trp Gly Gln Gly Thr Leu 100 105 110

Val Thr Val Ser Ser 115

<210> 96

<211> 105

<212> PRT

<213> Mus musculus

<400> 96

Gln Ile Val Leu Thr Gln Ser Pro Ala Ile Met Ser Ala Ser Leu Gly
1 5 10 15

Glu Glu Ile Thr Leu Thr Cys Ser Ala Ser Ser Ser Val Ser Tyr Met
20 25 30

His Trp Tyr Gln Gln Lys Ser Gly Thr Ser Pro Lys Leu Leu Ile Tyr 35 40 45

Arg Thr Ser Asn Leu Ala Ser Gly Val Pro Phe Arg Phe Ser Gly Ser 50 60

Gly Ser Gly Thr Phe Tyr Ser Leu Thr Ile Ser Ser Val Glu Ala Glu 65 70 75 80

Asp Ala Ala Asp Tyr Tyr Cys His Gln Trp Ser Met Tyr Thr Phe Gly
85 90 95

Gly Gly Thr Lys Leu Glu Ile Lys Arg 100 105

<210> 97

<211> 105

<212> PRT

<213> Artificial Sequence

-220-

<223> Description of Artificial Sequence: Synthetic humanized antibody

<400> 97

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Ser Ala Ser Ser Ser Val Ser Tyr Met 20 25 30

His Trp Tyr Gln Gln Lys Pro Gly Thr Ala Pro Lys Leu Leu Ile Tyr 35 40 45

Arg Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser Gly Ser 50 60

Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu 65 70 75 80

Asp Phe Ala Thr Tyr Tyr Cys His Gln Trp Ser Met Tyr Thr Phe Gly 85 90 95

Gln Gly Thr Lys Val Glu Ile Lys Arg 100 105

<210> 98

<211> 105

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
humanized antibody

<400> 98

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15

Glu Glu Val Thr Ile Thr Cys Ser Ala Ser Ser Ser Val Ser Tyr Met 20 25 30

His Trp Tyr Gln Gln Lys Pro Gly Thr Ala Pro Lys Leu Leu Ile Tyr

Arg Thr Ser Asn Leu Ala Ser Gly Val Pro Ser Arg Phe Ser Gly Ser

Gly Ser Gly Thr Phe Tyr Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu 65 70 75 80

Asp Phe Ala Thr Tyr Tyr Cys His Gln Trp Ser Met Tyr Thr Phe Gly
85 90 95

Gln Gly Thr Lys Val Glu Ile Lys Arg 100 105

<210> 99

<211> 105

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

humanized antibody

<400> 99

Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15

Glu Arg Ala Thr Ile Asn Cys Ser Ala Ser Ser Ser Val Ser Tyr Met 20 25 30

His Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr 35 40 45

Arg Thr Ser Asn Leu Ala Ser Gly Val Pro Asp Arg Phe Ser Gly Ser 50 60

Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala Glu 65 70 75 80

Asp Val Ala Val Tyr Tyr Cys His Gln Trp Ser Met Tyr Thr Phe Gly 85 90 95

Gln Gly Thr Lys Val Glu Ile Lys Arg 100 105